Run ora: October 5, 2005, 11:24:21; Search time 168 Seconds

(without alignments)

2115.671 Million cell updates/sec

Title: US-10-616-624-2

Perfect score: 4801

Sequence: 1 MALSWNEIRRKAIEFSKRWE.....QKMTSLLPPEKEIKKSKGKN 919

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 2105692 seqs, 386760381 residues

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : A\_Geneseq\_16Dec04:\*

1: geneseqp1980s:\*
2: geneseqp1990s:\*
3: geneseqp2000s:\*

4: geneseqp2001s:\*
5: geneseqp2002s:\*
6: geneseqp2003as:\*
7: geneseqp2003bs:\*

8: geneseqp2004s:\*

GenCore version 5.1.6

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OM protein - protein search, using sw model

Run on: October 5, 2005, 11:26:21; Search time 47 Seconds

(without alignments)

1881.345 Million cell updates/sec

Title: US-10-616-624-2

Perfect score: 4801

Sequence: 1 MALSWNEIRRKAIEFSKRWE.....QKMTSLLPPEKEIKKSKGKN 919

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 283416 segs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : PIR\_79:\*

1: pir1:\*

2: pir2:\*

3: pir3:\*

4: pir4:\*

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OM protein - protein search, using sw model

Run on: October 5, 2005, 11:24:51; Search time 183 Seconds

(without alignments)

2571.589 Million cell updates/sec

Title: US-10-616-624-2

Perfect score: 4801

Sequence: 1 MALSWNEIRRKAIEFSKRWE.....QKMTSLLPPEKEIKKSKGKN 919

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1612378 seqs, 512079187 residues

Total number of hits satisfying chosen parameters: 1612378

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : UniProt\_03:\*

1: uniprot\_sprot:\*
2: uniprot trembl::\*